

REPLACEMENT SHEET

1 MASTTPITMEDLQKALEAQSRALRAELAAG
 31 ASQSRRRPRPPRQRDSSTSGDDSGRDSGGPR
 61 RRRGNRGRGQRRDWSRAPPPPEERQESRSQ
 91 TPAPKP/SRAPPQQPOPPRMQTGRGGSAPRP
 121 ELGPPTNPFQAAVARGLRPPLHDPDTEAPT
 151 EACVTSWLWSEGQGA VFYRVDLHFTNLGTP
 181 PLDEDGRWDPALMYNPCGPEPPAHVVRAYN
 211 QPAGDVRGVWVGKGERTYAEQDFRVGGTRWH
 241 RLLRMPVRGLDGD SAPLPHTTERIETR SA
 271 RHPWRI RFGAPQAFLAGLLLATVAVG TARA
 301 GLQPRADMAAPPT LPQPPCAHGQH YGHHHH
 331 QLPFLGH DGH HGGTLRVGQH YRNASDVLP G
 361 HWLQGGWGCYNLSDWHQGTHVCHTKHMDFW
 391 CVEHARPPPATPTPLTTAANSTTAATPATA
 421 PAPCHAGLNDSCGGFLSGCGPMRLRHGADT
 451 RCGRLICGLSTTAQYPPTREFGCAMRWGLPP
 481 WELVVLTA RPE DGTGCRGVPAHPGARCPEL
 511 VSPMGRATCSPASALWLATANALSLDHALA
 541 AFVLSVPWVLIFMVCR RACRRRGAAAALTA
 571 VVLQGYNFPAYGEEAFTYLCTAPGCATQAP
 601 VPVRLAGVRFESKIVDGGCFAPWDLEATGA
 631 CICEIPTDVSCEGLGAWVPAAPCARIWNGT
 661 QRACTFWAVNAYSSGGYAQLASYFNPGGSY
 691 YKQYHPTACEVEPAFGHSDAACWGFPD TTV
 721 MSVFALASYVQHPHKTVRVKFHTETRTVWQ
 751 LSVAGVSCNVTT EHPFCNTPHGQLEVQVPP
 781 DPGDLVEYIMNYTGNQQSRWGLGSPNCHGP
 811 DWASPVCQRHSPDCSRLVGATPERPRLRLV
 841 DADDPLLR TAPGPGEVWVTPVIGSQARKCG
 871 LHIRAGPYGHATVEMPEWIIHAHTTSDPWHP
 901 PGPLGLKFKTVRPVALPRTLAPPRNV RV TG
 931 CYQCGTPALVEGLAPGGGNCHLT VNGEDVG
 961 AVPPGKFVTAALLNTPPPYQVSCGGESDRA
 991 SARVIDPAAQSFTGVVYGTHTTA VSETRQT
 1021 WAEWAAAHWWQLTLGATCALPLAGLLACCA
 1051 KCLYYLRGAIAPR

FIGURE 2